

## BACTERIA ASSOCIATED HOLOTHURIANS: THE KEY OF HABITAT PREFERENCE, DIET, AND FUNCTIONS

Ariani Hatmanti<sup>1)</sup> and Pradina Purwati<sup>1)</sup>

Research Center for Oceanography – Indonesian Institute of Science  
Jln. Pasir Putih I, Ancol Timur Jakarta Utara

### ABSTRACT

An investigation of bacteria as holothurians's feed carried out from May until June 2006 in Medane Bay and Kombal Bay, Lombok, Indonesia. The aims of this study are to know whether holothurians eat bacteria, to observe that bacteria are the key for holothurians' habitat preference, and to understand the functions of bacteria in holothurians intestine. The results showed there are many kind of bacteria in the substrates and inside intestine of sea cucumber. There are no specificity in performance, species and number of bacteria which found in anterior or posterior of intestine, as well in their substrates. Therefore, the holothurians do not choose specific bacteria for their feed. Bacteria are the consequence of substrates ingested. Some of them can be microflora in holothurians intestine. Bacteria inside holothurians produce some enzymes such as protease, amylase and agarase. This function was proven by the increasing of Total Organic Matters (TOM) in their feces. This evidence supporting Wiedmeyer (1992) statement that sea cucumber adds organic matters in the sediment. So this also firm holothurians' function in their habitat, as supplier for organic matters. The conclusion is that bacteria are eaten by holothurians by absorption, bacteria is not the key for microhabitat preference for holothurians, and inside the intestine of holothurians, digestion and degradation are the function of bacteria by producing their enzymes.

**Keywords:** *bacteria, holothurians, microhabitat preference, function*

### I. INTRODUCTION

For marine deposit - feeding invertebrates, the distribution of species with different life history strategies has long been known to be correlated with sediment organic concentration. Large populations of opportunistic species are found in sediments with enriched organic concentration, while equilibrium species populate low organic concentration sediments. Differences in both behavioral (e.g. feeding rate) and physiological (e.g. growth rate, reproductive output) adaptations determine the ability of species to establish populations in different environments. By systematically documenting differences in the way these factors vary as sediment

organic concentration varies for both opportunistic and equilibrium species, we can better understand the mechanisms underlying this correlation between sediment organic concentration and species distributions (Linton and Taghon, 2000).

Many publications (Massin, 1982; Michio *et al.*, 2003) explain that deposit-feeders holothurians eat bacteria, benthic microalgae, and organics matters associated with substrates. Bacteria can be feed for holothurians, they assumed to be absorbed in holothurians intestine. As we know, many animals feeding on sedimentary deposits are more nutritionally dependent on attached bacteria than on the nonliving organic debris (Mann, 1973). Sedimentary

bacteria are important in the functioning of benthic ecosystems, in that they are the primary remineralizers of organic matter. Deposit feeders are typically the dominant macrofaunal guild in soft sediments (Jumars, 1993), and ingest bacteria associated with sediments at a high frequency (Thayer, 1983; Wheatcroft *et al.*, 1990).

This research was done to answer the question that bacteria are one of the reasons why the holothurians choose their habitats. Generally, many animals feeding on sedimentary deposits are more nutritionally dependant on attached bacteria than on the nonliving organic debris. Bacteria and foraminifera may be major sources of food for atoll holothurians. The aims of this study are to know whether holothurians eat bacteria, to observe that bacteria

population is the key for holothurians' habitat preference, and to understand the functions of bacteria in holothurians intestine. The hypothesis was that bacteria are not the key for holothurians' habitat preference and they have some function inside holothurians.

## II. METHODS

This research was done in Kombal and Medane Bay, Lombok, Indonesia from May until June 2006. Kombal and Medane Bays are located in West of Lombok Island, 12 km northern from Senggigi. These bays position are  $116^{\circ}8'9.9''$  –  $116^{\circ}4'24''$  BT and  $8^{\circ}25'20.6''$  -  $8^{\circ}8'9.9''$  LS (Purwati *et al.*, 2008). Figure 1 below showed Lombok Island position in Indonesia archipelago map, while Kombal and Medane Bays location were showed in Figure 2.



Figure 1. Lombok Island Position (shown in red circle) in Indonesian Archipelago Map

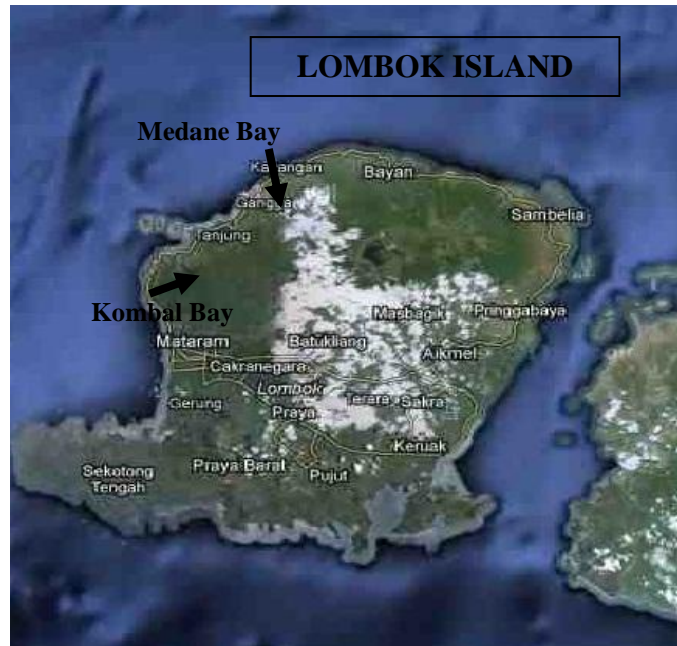


Figure 2. Kombal and Medane Bay Location in Lombok Island

Holothurians were collected at low tide, when the water going to high tide. The holothurians collected was then observed in the laboratory. Observation and description methods were used to analyses the relationship.

Samples were intestine of *Holothurian scabra*, *H. atra* and *H. leucospilota* and the substrates in front of their tentacles. The intestine was divided into anterior as fore part of the constriction (length was measured) and the posterior as 5 cm long from the posterior most. Both anterior and posterior of intestine were kept in a sterilized plastic bag, and then used for bacterial analysis. In order to provide selectivity information, sediment which was assumed to be fed was collected. A syringe was embedded 2-3 cm on the substrate in front of the mouth (in an area of 10 cm in diameter). Then the sediment was sucked out and the syringe was immediately taken off from the water, and the sediment was pushed into a sterile plastic bag. The samples were analyses by using each 1 gram of the samples (the anterior and posterior of

intestine and the substrate) were put in a test tube that fill by 9 ml sterile marine water for dilution from  $10^{-1}$  until  $10^{-4}$ . The  $10^{-3}$  and  $10^{-4}$  of dilution were put on a petridishes and pour with Nutrient Agar Media, by *pour plate* method. Those were kept in room temperature for 10 days. The bacteria growth on the NA media were counted by its performance (color, edge, shape, elevation), then purified on the same media. Pure bacteria were identified by Microgen<sup>TM</sup> Identification System. Pure bacteria are kept in  $-20^{\circ}\text{C}$  glycerol stock for culture collection and further analysis.

Skim Milk Agar (SMA) was used for protease producing bacteria analysis and Soluble Starch Agar (SSA) was used for amylase producing bacteria analysis. The samples were put on the media and kept for 3-5 days until they were analysis. Also pure isolates from Nutrient Agar were streaked on the each media for its analysis.

### III. RESULTS AND DISCUSSION

#### 3.1. Key for Habitat Preference and Diet

Bacteria can be feed for holothurians, they presumed to be absorbed in holothurians intestine. This study showed that some bacteria were

absorbed; it is proved by missing of some bacterial performance in their posterior intestine (Figure 3). Mostly of the bacteria amount decrease in the posterior intestine. It is need to be further analysis whether holothurians do not select bacteria for their feed by its species, but based on its composition of protein.

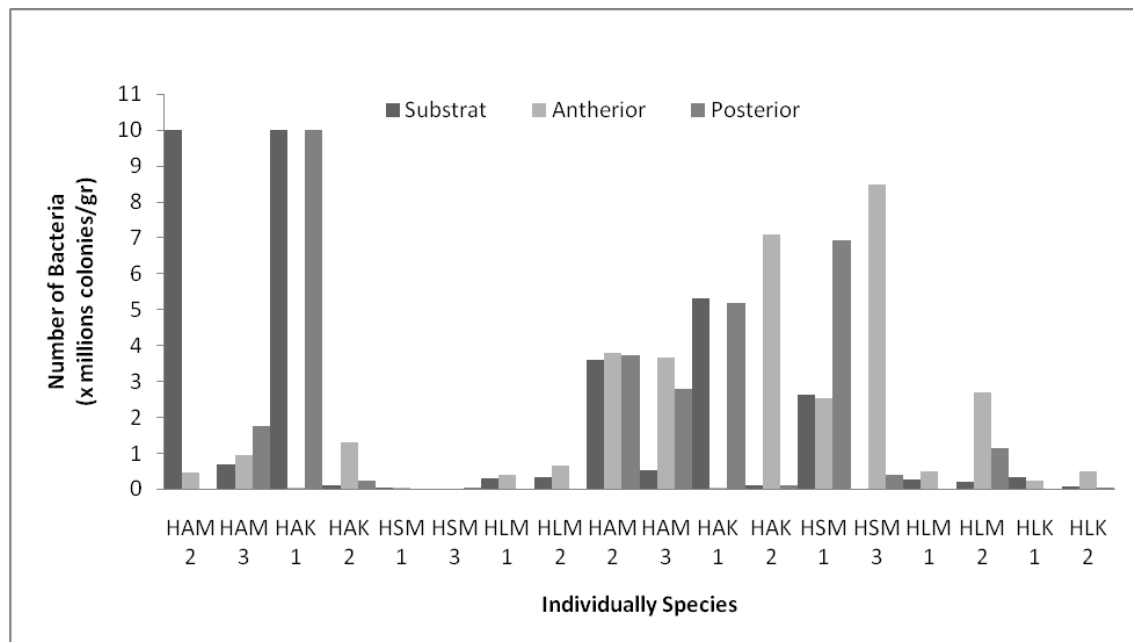


Figure 3. Comparison of number of cream colored, round glossy dome shaped bacteria between individually species in different locations : *Holothurian atra* Medana (HAM), *H. atra* Kombal (HAK), *H. scabra* Medana, *H. leucospilota* Medana (HLM), *H. leucospilota* Kombal (HLK)

Thus, the results showed that there are many kind of bacteria in the substrate and inside intestine of sea cucumber. There are no specificity in performance, species, and number of bacteria which found in anterior or posterior of sea cucumber intestine, as well in their substrates. Mean that holothurians do not choose specific bacteria for their feed. This phenomenon results a conclusion that bacteria are not the reason why the holothurians choose a habitat. This microorganism is not the key for microhabitat preference. Bacteria are consequence of substrates ingested.

The results showed that there are a lot of variations on bacterial morphology shown on the plate. But there are unequivocal pattern among substrate, anterior and posterior. This can be shown from the data sorting of each holothurians. In substrate, bacteria more varied than those in intestine, especially for *H. leucospilota*. Assumed these organisms choose richer habitat than *H. atra* and *H. scabra*. Based on that, variation of bacteria morphological performance in anterior less varied than those in substrate. Holothurians did not select bacteria they ingest from substrate. It can be shown that individually

holothurians from the same species ate difference bacteria, as shown in their anterior intestine. It is that holothurians filter randomized bacteria for them to eat. Purwati *et al.* (2006) in their report declared that from Friedman test there are no preferences of selectivity to particle size of sediment for *H. atra*, but *H. scabra* prefer to soft than hard particles. It is also stated that there is no selectivity of diatoms for their diet.

There is one morphological performance which always be found in holothurians substrate and intestine, such as cream colored with round glossy dome shaped. Figure 3 shows that there are no special pattern of amount of the bacteria, both in substrate and intestine. It is presumed that these bacteria have no meaning to holothurians diets.

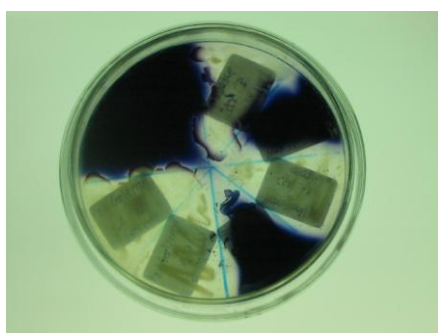
### 3.2. Functions of Bacteria in Holothurians

Bacteria in association with holothurians have many functions as well digestion and degradation of its feed, because they produce some enzymes such as protease, amylase and agarase. Some of them can be microflora in holothurians intestine. There are two functions of bacteria inside sea cucumber, that is bacteria absorbed by holothurians as feed, and bacteria which have the ability to degrade organic matters inside their intestine. The

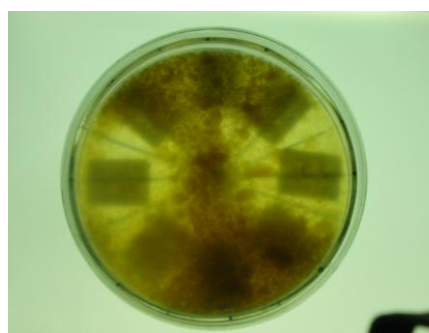
function is related with increasing of Total Organic Matters (TOM) in their fecal pellet. Wiedmeyer (1992) stated that sea cucumber adds organic matters in their environment. Sea cucumbers turn their less nutrient substrate into rich organic matter substrate. It is one of holothurian's benefits to their environment. So this also firm holothurians' function in their habitat, as supplier for organic matters.

Nutrient-degrader bacteria in holothurians's intestine have been proved in this experiment, as well starch and protein degraders. The aim of this experiment was to prove that bacteria are not only their feed, but also have some functions, related to their enzyme. Starch and protein have been selected for substrate tested because they are nutrient sources in great quantities in the sea.

The data show that in *H. atra* 45.5% bacteria degrade protein (produce protease) and 30.3% degrade starch (amylase producing bacteria). In *H. scabra* 73.3% bacteria produce protease but only 13.3% produce amylase, while in *H.leucosphilota* 73.7% are protease producing bacteria and only 26.3% bacteria produce amylase. Variation percentage between species of holothurians was depending on their substrate composition. Figure 4 shows the ability of bacteria to degrade starch (a) and protein (b).



(a)



(b)

Figure 4. Picture of starch degrading bacteria (a) and protein degrading bacteria (b).

Figure 4 (a) point out Starch Soluble Agar (SSA+50% seawater) which have been grown by bacteria. Clear zone shows that the bacteria produce amylase. Figure 4 (b) shows Skim Milk Agar (SMA+50% seawater) which have been grown by bacteria. The clear zone refers to protease producing bacteria.

Degrading bacteria were found in sea cucumber intestine with pattern as follows: in *H. atra* those can be found in substrate, anterior and posterior intestine, but mostly in posterior. In *H. scabra* degrading bacteria can be found in substrate, anterior and posterior. In *H. leucosphilota* amylase and protease producing bacteria were found mostly in substrate. There were two possibility that can be presumed from this statement, first that *H. leucosphilota* prefer bacterial rich substrate, or the second, degrading bacteria compose substrate become organic matter rich substrate which *H. leucosphilota* would like to use for their feed and protection.

For addition, in anterior of *H. leucosphilota*'s intestine from Medane Bay, there always be found agar lysis bacteria, even in posterior it can be liquidized agar. The bacteria supposed to have agarase. Assumption that *H. leucosphilota*'s substrate was composed of agar like compound as well as karaginan and gelatin can be one of reason the presence of agarase producing bacteria in their intestine. It is strengthen that inside sea cucumber there are substrate-degrading bacteria.

Deposit feeders have been shown to affect bacterial densities (Moriarty *et al.*, 1985; Grossmann & Reichardt, 1991), growth (Deming & Colwell, 1982; Plante *et al.*, 1989), activity (Juniper, 1981; Deming & Colwell, 1982; Aller & Yingst, 1985), and impact community compositions (Dobbs & Guckert, 1988; Duchene *et al.*, 1988; Ward-Rainey *et al.*,

1996). Historically, the relationship between deposit feeders and sedimentary bacteria has been studied in the context of a classic predator-prey (+, -) interaction. We argue here that treatment of deposit-feeder ingestion as a disturbance is more appropriate. Most importantly, it is not clear that digestion of bacteria provides a net gain to deposit feeders (Lopez & Levinton, 1987; Kemp, 1990). When compared to the typical relationship of a predator and prey, the effect of a deposit feeder on ingested microbes occurs over larger spatial scales, influences communities as opposed to individuals, and is less selective. Moreover, a disturbance disrupts the physical environment as well as biota. Deposit feeding alters geochemical gradients within ingested sediments (Aller, 1982; Plante & Jumars, 1992), dramatically affects the quantity and architecture of microbial biofilms (Decho & Lopez, 1993), and can affect grain size, sorting, water content and compaction of sediments (Gray, 1974; Rhoads & Boyer, 1982).

#### IV. CONCLUSION

The bacteria eaten by holothurians were by absorption process, bacteria were not the key for microhabitat preference for holothurians, and inside the intestine of holothurians, digestion and degradation were the function of bacteria by producing their enzymes.

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